## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	_/0/533,3 <i>0</i> 0
Source:	P4/10
Date Processed by STIC:	5/12/05

## ENTERED



PCT

RAW SEQUENCE LISTING DATE: 05/12/2005
PATENT APPLICATION: US/10/533,300 TIME: 08:22:30

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\05122005\J533300.raw

```
3 <110> APPLICANT: Meder, Wolfgang
              Wendland, Martin
      4
              John, Harald
      5
              Richter, Rudolf
      7
              Meyer, Markus
              Forssmann, Wolf-Georg
     10 <120> TITLE OF INVENTION: Human Chondroosteomodulin (TIG2), Production, and Use for
the
              Treatment or Diagnosis of Bone Diseases, Cartilage Diseases,
     11
              Obesity, Inflammatory Diseases, and Skin Diseases
     12
     14 <130> FILE REFERENCE: 2388.0010000/SRL/BLS
C--> 16 <140> CURRENT APPLICATION NUMBER: US/10/533,300
C--> 16 <141> CURRENT FILING DATE: 2005-04-29
     16 <150> PRIOR APPLICATION NUMBER: PCT/EP03/11799
     17 <151> PRIOR FILING DATE: 2003-10-24
     19 <160> NUMBER OF SEQ ID NOS: 16
     21 <170> SOFTWARE: PatentIn version 3.3
     23 <210> SEQ ID NO: 1
     24 <211> LENGTH: 134
     25 <212> TYPE: PRT
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                                        25
     38 Glu Ser Ala Val Asp Thr Pro Phe Pro Ala Gly Ile Phe Val Arg Leu
     42 Glu Phe Lys Leu Gln Gln Thr Ser Cys Arg Lys Arg Asp Trp Lys Lys
     46 Pro Glu Cys Lys Val Arg Pro Asn Gly Arg Lys Arg Lys Cys Leu Ala
     50 Cys Ile Lys Leu Gly Ser Glu Asp Lys Val Leu Gly Arg Leu Val His
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                                             90
     54 Cys Pro Ile Glu Thr Gln Val Leu Arg Glu Ala Glu Glu His Gln Glu
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     67 <211> LENGTH: 1186
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71 <220> FEATURE: 72 <223> OTHER INFORMATION: Description of artificial sequence: GORI-28 cDNA 74 <400> SEQUENCE: 2 75 tggtccctgt cttctcttgc agagaatgga ggatgaagat tacaacactt ccatcagtta 77 cqqtqatqaa taccctqatt atttagactc cattqtggtt ttggaggact tatccccctt 120 79 ggaagccagg gtgaccagga tetteetggt ggtggtetae ageategtet getteetegg 180 81 gattctgggc aatggtctgg tgatcatcat tgccaccttc aagatgaaga agacagtgaa 240 83 catggtctgg ttcctcaacc tggcagtggc agatttcctg ttcaacgtct tcctcccaat 300 85 ccatatcacc tatgccgcca tggactacca ctgggttttc gggacagcca tgtgcaagat 360 87 cagcaacttc cttctcatcc acaacatgtt caccagcgtc ttcctgctga ccatcatcag 420 89 ctctgaccgc tgcatctctg tgctcctccc tgtctggtcc cagaaccacc gcagcgttcg 480 91 cctggcttac atggcctgca tggtcatctg ggtcctggct ttcttcttga gttccccatc 540 93 totogtotto ogggacacag ocaacotgoa tgggaaaata tootgottoa acaacottoag 600 95 cctgtccaca cctgggtctt cctcgtggcc cactcactcc caaatggacc ctgtggggta 660 97 tageeggeae atggtggtga etgteaceeg etteetetgt ggetteetgg teecagteet 720 99 catcatcaca gettgetace teacéategt gtgeaaactg cagegeaace geetggeeaa 780 101 gaccaagaag cccttcaaga ttattgtgac catcatcatt accttcttcc tctgctggtg 840 103 cccctaccac acactcaacc tcctagaget ccaccacact gecatgeetg getetgtett 105 cagectgggt ttgecectgg ceactgeect tgecattgee aacagetgea tgaaceceat 960 107 tetqtatqtt tteatqqqte aggaetteaa qaagtteaag gtggeeetet tetetegeet 1020 109 ggtcaatget ctaagtgaag atacaggeca ctetteetac cecagecata gaagetttac 1080 111 caaqatgtca tcaatgaatg agaggacttc tatgaatgag agggagaccg gcatgctttg 1140 113 atcctcactg tggaacccct caatggactc tctcaaccca gggaca 1186 116 <210> SEQ ID NO: 3 117 <211> LENGTH: 18 118 <212> TYPE: DNA 119 <213> ORGANISM: Artificial Sequence 121 <220> FEATURE: 122 <223> OTHER INFORMATION: Description of artificial sequence: Primer TIG2oli1 124 <400> SEQUENCE: 3 125 gccagggtga cacggaag 1.8 128 <210> SEQ ID NO: 4 129 <211> LENGTH: 17 130 <212> TYPE: DNA 131 <213> ORGANISM: Artificial Sequence 133 <220> FEATURE: 134 <223> OTHER INFORMATION: Description of artificial sequence: Primer TIG2oli2 136 <400> SEQUENCE: 4 137 gaggcaccac gcagctc 17 140 <210> SEQ ID NO: 5 141 <211> LENGTH: 33 142 <212> TYPE: DNA 143 <213> ORGANISM: Artificial Sequence 145 <220> FEATURE: 146 <223> OTHER INFORMATION: Description of artificial sequence: Primer COMoli1 148 <400> SEQUENCE: 5 149 ctctcqaqaa aagagagctc acggaagccc agc 33 152 <210> SEO ID NO: 6 153 <211> LENGTH: 34

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- 154 <212> TYPE: DNA
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- 157 <220> FEATURE:
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- 164 <210> SEQ ID NO: 7
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- 166 <212> TYPE: DNA
- 167 <213> ORGANISM: Artificial Sequence
- 169 <220> FEATURE:
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- 172 <400> SEQUENCE: 7
- 173 ggccatgtgc aagatcagca act
- 176 <210> SEQ ID NO: 8
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- 188 <210> SEQ ID NO: 9
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- 191 <213> ORGANISM: Artificial Sequence
- 193 <220> FEATURE:
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- 212 <210> SEQ ID NO: 11
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- 214 <212> TYPE: DNA
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- 217 <220> FEATURE:
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- 221 cagacaacat aacggtgaat ga
- 224 <210> SEQ ID NO: 12
- 225 <211> LENGTH: 19
- 226 <212> TYPE: DNA

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- 251 <213> ORGANISM: Artificial Sequence
- 253 <220> FEATURE:
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- 256 <400> SEQUENCE: 14
- 257 ctgtagacca ccaccaggaa ga
- 260 <210> SEQ ID NO: 15
- 261 <211> LENGTH: 20
- 262 <212> TYPE: DNA
- 263 <213> ORGANISM: Artificial Sequence
- 265 <220> FEATURE:
- 266 <223> OTHER INFORMATION: Description of artificial sequence: Primer GORI280111
- 268 <400> SEQUENCE: 15
- 272 <210> SEQ ID NO: 16
- 273 <211> LENGTH: 20
- 274 <212> TYPE: DNA
- 275 <213> ORGANISM: Artificial Sequence
- 277 <220> FEATURE:
- 278 <223> OTHER INFORMATION: Description of artificial sequence: Primer GORI2801i2
- 280 <400> SEQUENCE: 16
- 281 tgtccctggg ttgagagagt 20

VERIFICATION SUMMARY

DATE: 05/12/2005

PATENT APPLICATION: US/10/533,300

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